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09/647,978

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**OCTOBER 6, 2000** 

FOR:

GENE ENCODING SYNTAXIN INTERACTING PROTEIN

## SEQUENCE LISTING CERTIFICATION

I hereby certify that the paper copy of the Sequence Listing is identical to the computer readable copy of the Sequence Listing and that the Sequence Listing does not include matter which goes beyond the disclosure in the International Application as filed.

Date

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Registration No. 27,610

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<110> Warner-Lambert Company

<120> Gene Encoding Syntaxin Interacting Profein

<130> 09/647,978

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<151> 1999-04-19

<150> 60/082,454

<151> 1998-04-20

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M

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35 40 45

Ile His Glu Val Ile Pro Gly Gly Asp Cys Tyr Lys Asp Gly Arg Leu
50 55 60

Lys Pro Gly Asp Gln Leu Val Ser Ile Asn Lys Glu Ser Met Ile Gly 65 70 75 80

Val Ser Phe Glu Glu Ala Lys Ser Ile Ile Thr Arg Ala Lys Leu Arg
85 90 95

Ser Glu Ser Pro Trp Glu Me Ala Phe Ile Arg Gln Lys Ser Tyr Cys
100 105 110

Gly His Pro Gly Asn Ile Cys Cys Pro Ser Pro Gln Val Ser Glu Asp 115 120 125

Cys Gly Pro Gln Thr Ser Thr Phe Thr Leu Leu Ser Ser Pro Ser Glu
130 135 140

Thr Leu Leu Pro Lys Thr Ser Ser Thr Pro Gln Thr Gln Asp Ser Thr 145 150 155 160

Phe Pro Ser Cys Lys Ala Ile Gln Thr Lys Pro Glu His Asp Lys Thr
165 170 175

Glu	His	Ser	Pro 180	Ile	Thr	Ser	Leu	Asp 185	Asn	Ser	Pro	Ala	Asp 190	Thr	Ser
Asn	Ala	Asp 195	Ile	Ala	Pro	Ala	Trp 200	Thr	Asp	Asp	Asp	Ser 205	Gly	pro	Gln
Gly	Lys 210	Ile	Ser	Leu	Asn	Pro 215	Ser	Val	Arg	Leu	Lys 220	Ala	€1u	Lys	Leu
Glu 225	Met	Ala	Leu	Asn	Tyr 230	Leu	Gly	Ile	Gln	Pro 235	Thr	Lys	Glu	Gln	Arg 240
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Gln	Leu 290	Leu	Pro	Cys	Asp	Ser 295	Leu	<b>El</b> u	Ala	Asp	Glu 300	Val	Gly	Lys	Leu
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Val 385	Ser	Glu	Leu	Lys	Ala 390	Gln	Leu	Ala	Asp	Tyr 395	Ser	Asp	Gln	Asn	Lys 400
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<213> Human

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Ile Gln Pro Glu Asn Ley Ser Cys Thr Ser Leu Ile Glu Ala Ser Gly
115 120 125

Glu Tyr Gly Pro Gln/Ala Ser Thr Leu Ser Leu Phe Ser Ser Pro Pro 130 135 140

Glu Ile Leu Ile ro Lys Thr Ser Ser Thr Pro Lys Thr Asn Asn Asp 145 150 155 160

Ile Leu Ser Ser Cys Glu Ile Lys Thr Gly Tyr Asn Lys Thr Val Gln
165 170 175

Ile Pro Ile Thr Ser Glu Asn Ser Thr Val Gly Leu Ser Asn Thr Asp

al

Val Ala Ser Ala Trp Thr Glu Asn Tyr Gly Leu Gln Glu Lys I e Ser 195 200 205

185

Leu Asn Pro Ser Val Arg Phe Lys Ala Glu Lys Leu Glu Met Ala Leu 210 215 220

Asn Tyr Leu Gly Ile Gln Pro Thr Lys Glu Gln His Gln Ala Leu Arg 225 230 235 240

Gln Gln Val Gln Ala Asp Ser Lys Gly Thr Val Ser Phe Gly Asp Phe
245 250 255

Val Gln Val Ala Arg Asn Leu Phe Cys Leu Gln Leu Asp Glu Val Asn 260 265 270

Val Gly Ala His Glu Ile Ser Asn Ile Leu Asp Ser Gln Leu Leu Pro 275 280 285

Cys Asp Ser Ser Glu Ala Asp Glu Met Glu Arg Leu Lys Cys Glu Arg 290 295 300

Asp Asp Ala Leu Lys Glu Val Asn Thr Leu Lys Glu Lys Leu Leu Glu 305 310 315 320

Ser Asp Lys Gln Arg Lys Gln Leu Thr Glu Glu Leu Gln Asn Val Lys 325 330 335

Gln Glu Ala Lys Ala Val Val Glu Glu Thr Arg Ala Leu Arg Ser Arg
340 345 350

Ile His Leu Ala Glu Ala Gln Arg Gln Ala His Gly Met Glu Met
355 360 365

Asp Tyr Glu Glu Val Ile Arg Leu Leu Glu Ala Lys Ile Thr Glu Leu 370 375 380

Lys Ala Gln Ley Ala Asp Tyr Ser Asp Gln Asn Lys Glu Ser Val Gln 385 390 395 400

Asp Leu Lys Lys Arg Ile Met Val Leu Asp Cys Gln Leu Arg Lys Ser

Glu Met Ala Arg Lys Thr Phe Glu Ala Ser Thr Glu Lys Leu Leu His
420 425 430

Phe Va / Glu Ala Ile Gln Glu Val Phe Ser Asp Asn Ser Thr Pro Leu

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